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RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/876,252

DATE: 12/07/2001
TIME: 16:39:27

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3 <110> APPLICANT: Behan, Dominic P.
4 Lehmann-Bruinsma, Karin
5 Chalmers, Derek T.
6 Lowitz, Kevin P.
7 Lin, I-Lin
8 Dang, Huong T.
9 Chen, Ruoping
10 Liaw, Chen W.
12 <120> TITLE OF INVENTION: Non-Endogenous Constitutively Activated Human G Protein
Coupled Receptors
14 <130> FILE REFERENCE: AREN-0054
16 <140> CURRENT APPLICATION NUMBER: 09/876,252
17 <141> CURRENT FILING DATE: 2001-06-07
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 123 agtccattgc ttagatatacg ttttggaaacc atggctccca ctgggtttagg ttcccttgacc 180
 125 gtgaatagta cagctgtgcc cacaacacca gcagcattta agagcctaaa ctgcctt 240
 127 cagatcaccc tttctgttat aatgtatattc attctgtttg tgtctttct tggaaacttg 300
 129 gttgtttgcc tcattgttta ccaaaaagct gccatgaggt ctgcaatcaa catccctt 360
 131 gcccggcttag ctttgcaga catgttgctt gcagtgtga acatgccctt tgccctggta 420
 133 actattctta ctaccggatg gatTTTtggg aaattcttct gttaggtatc tgctatgttt 480
 135 ttctggttat ttgtataga aggatgtacc atcctgctca tcatttagat agataggttc 540
 137 cttattatacg tccagaggca ggataagcta aaccatata gagctaagg tctgattgca 600
 139 gtttcttggg caacttcctt ttgtgttagt tttcctttag ccgttagggaa ccccgacctg 660
 141 cagataccctt cccgagctcc ccagtgtgtg tttgggtaca caaccaatcc aggctaccag 720
 143 gcttatgtga ttttggatttc tctcattttct ttcttcatac cttccctgtt aatactgtac 780
 145 tcattttatgg gcataactcaa cacccttccg cacaatgcct tgaggatcca tagtaccct 840
 147 gaaggatatacgccca ggccagcaa ctgggtctca tgagtctgca gagacccctt 900
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 157 gttgcctgg acatgatgcc taagtccttc aagttttgc cgccagctccc tggcacaca 1200
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 175 Pro Pro Phe Gln His Pro Asp Leu Ser Pro Leu Leu Arg Tyr Ser Phe
 176 35 40 45
 178 Glu Thr Met Ala Pro Thr Gly Leu Ser Ser Leu Thr Val Asn Ser Thr
 179 50 55 60
 181 Ala Val Pro Thr Thr Pro Ala Ala Phe Lys Ser Leu Asn Leu Pro Leu
 182 65 70 75 80
 184 Gln Ile Thr Leu Ser Ala Ile Met Ile Phe Ile Leu Phe Val Ser Phe
 185 85 90 95
 187 Leu Gly Asn Leu Val Val Cys Leu Met Val Tyr Gln Lys Ala Ala Met
 188 100 105 110
 190 Arg Ser Ala Ile Asn Ile Leu Leu Ala Ser Leu Ala Phe Ala Asp Met
 191 115 120 125
 193 Leu Leu Ala Val Leu Asn Met Pro Phe Ala Leu Val Thr Ile Leu Thr
 194 130 135 140
 196 Thr Arg Trp Ile Phe Gly Lys Phe Phe Cys Arg Val Ser Ala Met Phe
 197 145 150 155 160
 199 Phe Trp Leu Phe Val Ile Glu Gly Val Ala Ile Leu Leu Ile Ile Ser
 200 165 170 175
 202 Ile Asp Arg Phe Leu Ile Ile Val Gln Arg Gln Asp Lys Leu Asn Pro
 203 180 185 190
 205 Tyr Arg Ala Lys Val Leu Ile Ala Val Ser Trp Ala Thr Ser Phe Cys
 206 195 200 205
 208 Val Ala Phe Pro Leu Ala Val Gly Asn Pro Asp Leu Gln Ile Pro Ser
 209 210 215 220
 211 Arg Ala Pro Gln Cys Val Phe Gly Tyr Thr Thr Asn Pro Gly Tyr Gln
 212 225 230 235 240
 214 Ala Tyr Val Ile Leu Ile Ser Leu Ile Ser Phe Phe Ile Pro Phe Leu
 215 245 250 255
 217 Val Ile Leu Tyr Ser Phe Met Gly Ile Leu Asn Thr Leu Arg His Asn
 218 260 265 270
 220 Ala Leu Arg Ile His Ser Tyr Pro Glu Gly Ile Cys Leu Ser Gln Ala
 221 275 280 285
 223 Ser Lys Leu Gly Leu Met Ser Leu Gln Arg Pro Phe Gln Met Ser Ile
 224 290 295 300
 226 Asp Met Gly Phe Lys Thr Arg Ala Phe Thr Thr Ile Leu Ile Leu Phe
 227 305 310 315 320

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 230 325 330 335
 232 Ala Thr Phe Ser Lys His Phe Tyr Tyr Gln His Asn Phe Phe Glu Ile
 233 340 345 350
 235 Ser Thr Trp Leu Leu Trp Leu Cys Tyr Leu Lys Ser Ala Leu Asn Pro
 236 355 360 365
 238 Leu Ile Tyr Tyr Trp Arg Ile Lys Lys Phe His Asp Ala Cys Leu Asp
 239 370 375 380
 241 Met Met Pro Lys Ser Phe Lys Phe Leu Pro Gln Leu Pro Gly His Thr
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 245 405 410 415
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 260 gcgctagccc tctgggtctt cctgcgcgcg ctgcgcgtgc actcgggtgg gacgtgtac 180
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 266 ttccagatga acatgtacgg cagctgcata ttccgtatgc tcatcaacgt ggaccgctac 360
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 270 ctctgcctgg gcgtgtggc gtcatccctg gtgtttcccg tgcccgccgc cccgtgcac 480
 272 aggcctctcg gttgcgccta cccggacccc gaggtgcgc tatgttcgaa gagtttcagc 540
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 276 ctgtgcctcc tggcgccgg ggtctactcg tcggccgag tttctggac gctggcgcc 660
 278 cccgacgcac cgcagagcca gccggcgccgg aagaccgtgc gcctccctgt ggctaaccctc 720
 280 gtcatcttcc tgctgtgtt cgtgcctac aacagcacgc tggcggtcta cggctgtctg 780
 282 cggagcaagc tggtgccggc cagctgtgcct gcccgcata gctgtgcgcgg ggtgtatgc 840
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 308 Leu Ala Ala Gly Leu Pro Leu Asn Ala Leu Ala Leu Trp Val Phe Leu
 309 35 40 45
 311 Arg Ala Leu Arg Val His Ser Val Val Ser Val Tyr Met Cys Asn Leu

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318					85				90							95
320	Thr	Gly	Ala	Ile	Phe	Gln	Met	Asn	Met	Tyr	Gly	Ser	Cys	Ile	Phe	Leu
321					100				105							110
323	Met	Leu	Ile	Asn	Val	Asp	Arg	Tyr	Ala	Ala	Ile	Val	His	Pro	Leu	Arg
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326	Leu	Arg	His	Leu	Arg	Arg	Pro	Arg	Val	Ala	Arg	Leu	Leu	Cys	Leu	Gly
327					130				135							140
329	Val	Trp	Ala	Leu	Ile	Leu	Val	Phe	Ala	Val	Pro	Ala	Ala	Arg	Val	His
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332	Arg	Pro	Ser	Arg	Cys	Arg	Tyr	Arg	Asp	Leu	Glu	Val	Arg	Leu	Cys	Phe
333					165				170							175
335	Glu	Ser	Phe	Ser	Asp	Glu	Leu	Trp	Lys	Gly	Arg	Leu	Leu	Pro	Leu	Val
336					180				185							190
338	Leu	Leu	Ala	Glu	Ala	Leu	Gly	Phe	Leu	Leu	Pro	Leu	Ala	Ala	Val	Val
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341	Tyr	Ser	Ser	Gly	Arg	Val	Phe	Trp	Thr	Leu	Ala	Arg	Pro	Asp	Ala	Thr
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344	Gln	Ser	Gln	Arg	Arg	Arg	Lys	Thr	Val	Arg	Leu	Leu	Leu	Ala	Asn	Leu
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353	Asp	Arg	Val	Arg	Gly	Val	Leu	Met	Val	Met	Val	Leu	Leu	Ala	Gly	Ala
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356	Asn	Cys	Val	Leu	Asp	Pro	Leu	Val	Tyr	Tyr	Phe	Ser	Ala	Glu	Gly	Phe
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